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of breed,

Genome

File 🔽

☐ 1: H02533. yj40fl1.rl Soares...[gi:865466]

Links

Details

Boo

### **IDENTIFIERS**

dbEST Id:

262413

EST name: GenBank Acc: yj40f11.r1 H02533

GenBank gi:

865466

GDB Id:

563320

# CLONE INFO

Clone Id:

IMAGE: 151245 (5')

Insert length:
DNA type:

739 cDNA

## **PRIMERS**

Sequencing:

M13RP1

PolyA Tail:

Unknown

## SEQUENCE

TTTTACTGCACAATTCACAAGCATGTTTTCCTGGTGAATTGGACTGAAAATTACATTTTG
ACAACTTTTTTTCCTTTTATCCCCAACTTTTTGCCAGAAAGCAGAAAAATGCTCTATTTTT
ATAAAGAAAGATTAAATTCTCCCAATGATATTTTAAAAAAATATCAACCTACATGCACTTTA
GAATGTAAAATAACAATGACTATTTTAAACTCGAAGACCCACTATTTTGAGTATTTTTA
TAGACTTTAAATACTGGGTTTTTTTCCTCCTTCAATCTCAGGGCTTTTCTCCATCTTTTA
AGGCAGCCTCTGTAACTCCCTTTTGTCCATAGGTGTTGCGTGCCTCTGCATCTGTGGGGG
AAGTATTATTTAAATAAAATTTATGTTACAGGGGATAACTTTATTTTAAAGGGGCNGGGGC

TTGTCTATACACAAGGGTATGTATG

Quality:

High quality sequence stops at base: 331

Entry Created: Last Updated: Jun 20 1995 Jun 20 1995

## COMMENTS

Insert Size: 739

High qality sequence stops: 331 Source: IMAGE Consortium, LLNL

This clone is available royalty-free through LLNL; contact

the IMAGE Consortium (info@image.llnl.gov) for further

information.

## LIBRARY

Lib Name:

Soares placenta Nb2HP

Organism:

Homo sapiens

Sex:

Female

Not I

Organ:

placenta

Lab host:

Develop. stage: placenta obtained at birth (full term)

Vector:

DH10B (ampicillin resistant)

R. Site 1:

pT7T3D (Pharmacia) with a modified polylinker

R. Site 2:

Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer double-stranded cDNA was ligated to Eco RI adaptors

(Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of the modified pT7T3 vector. Library went through one round of normalization. Library constructed by

Bento Soares and M. Fatima Bonaldo.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel:

314 286 1800 314 286 1810

Fax:

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

The WashU-Merck EST Project

Authors:

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M. , Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

Wilson, R.

Year:

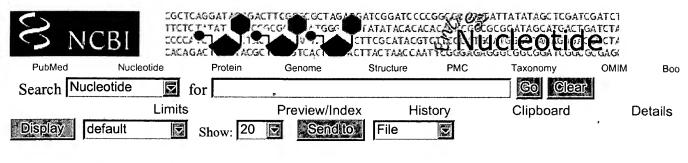
1995

Status:

Unpublished

MAP DATA

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☐ 1: AA010665. ze19f06.s1 Soares...[gi:1471692]

Links

## IDENTIFIERS

 dbEST Id:
 621531

 EST name:
 ze19f06.s1

 GenBank Acc:
 AA010665

 GenBank gi:
 1471692

 GDB Id:
 1276003

CLONE INFO

Clone Id: IMAGE:359459 (3')
Source: IMAGE Consortium, LLNL

Insert length: 495 DNA type: cDNA

PRIMERS

Sequencing: mob.REGA+ET PolyA Tail: Unknown

SEQUENCE

Entry Created: Jul 16 1996 Last Updated: Nov 29 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (<u>info@image.llnl.gov</u>) for further

information.

LIBRARY

Lib Name: Soares\_fetal\_heart\_NbHH19W

Organism: Homo sapiens
Sex: unknown
Organ: heart
Develop. stage: 19 weeks

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: Not I
R. Site 2: Eco RI

Description: 1st strand cDNA was primed with a Not I - oligo(dT) primer

adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector (Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by M.Fatima Bonaldo. This library was constructed from the same fetus as the fetal lung library, Soares fetal lung NbHL19W.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax: 314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

The WashU-Merck EST Project

Authors:

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
Holman,M., Hultman,M., Kucaba,T., Le,M., Lennon,G., Marra,M.
, Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,

Wilson, R.

Year:

1995

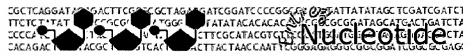
Status:

Unpublished

MAP DATA

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Structure

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default  $\overline{\Box}$ 

Nucleotide

Show: 20

File  $\nabla$ 

☐ 1: N69022. za71f10.s1 Soares...[gi:1225183]

Limits

Links

### IDENTIFIERS

dbEST Id:

475103

EST name: GenBank Acc:

za71f10.s1 N69022

GenBank gi:

1225183

GDB Id:

1242949

## CLONE INFO

Clone Id:

IMAGE:298027 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

**CDNA** 

### **PRIMERS**

Sequencing:

m13 -40 forward

PolyA Tail:

Unknown

# SEQUENCE

CTTTCCATCTATAATATGTGCTACAAATATATTTTTCAAAGGTCCAACCCAGGTTAGGAG GCTTCAAGGACCCTTTCTTAGCTACTGATTTTAGTAATTAAAAAAATAAACACAATAAAC ACCCCTGGCATCTTTGTGAGCGCCCCCCTGCGGAGAGCATCGAGGGCTGTCAGGCATCTT CCTCCTGCAGCCGCAGCCAATGGAGCACACGGAGAACTGGAGTCAGGTGATCACGAAGGG GCTTATTTCACTGGGCGTGGTCTACGTTGCCACATGTGTCGCTTTGGGATGAATGCATTT TTTNCAATGTGCTGCTTACACAATTATGCTGGTTCCAAAAACACAGTTGGACTTGGACAAT CTAACATAGGATTTTAATCCTTAGAGGT

Quality:

High quality sequence stops at base: 307

Entry Created: Last Updated:

Mar 13 1996 Mar 13 1996

# COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further information.

# LIBRARY

Lib Name:

Soares fetal lung NbHL19W

Organism:

Homo sapiens

Organ:

lung

Develop. stage: 19 weeks

Lab host:

DH10B (ampicillin resistant)

Vector:

pT7T3D (Pharmacia) with a modified polylinker

R. Site 1: R. Site 2:

Not I Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer double-stranded cDNA was size selected, ligated to Eco RI

adapters (Pharmacia), digested with Not I and cloned into

the Not I and Eco RI sites of a modified pT7T3 vector

(Pharmacia). Library went through one round of normalization to a Cot = 5. Library constructed by Bento Soares and

M.Fatima Bonaldo. This library was constructed from the same

fetus as the fetal heart library, Soares fetal heart

NbHH19W.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax: 314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

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The WashU-Merck EST Project

Authors:

Hillier,L., Clark,N., Dubuque,T., Elliston,K., Hawkins,M.,
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Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,

Wilson, R.

Year:

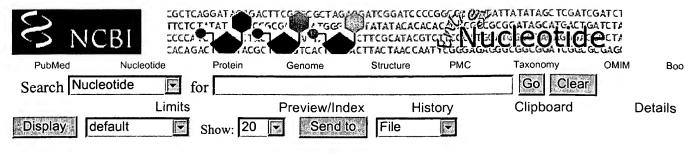
1995

Status:

Unpublished

### MAP DATA

<u>Disclaimer</u> | <u>Write to the Help Desk</u> <u>NCBI</u> | <u>NLM</u> | <u>NIH</u>



☐ 1: D60118. HUM087A05A Clonte...[gi:961757]

Links

## IDENTIFIERS

dbEST Id: 335105 EST name: HUM087A05A GenBank Acc: D60118 GenBank gi: 961757

CLONE INFO

GEN-087A05 (3') Clone Id: **CDNA** 

DNA type:

**PRIMERS** 

PolyA Tail: Unknown

SEQUENCE

AAAACTGAGGGAACTCATGCTTTAATAGACACTGAAAATCACAAAGGRGGAAGGCCAAGT GCCTTAGCAATCTCAATAAAAATATGAVGTTCTTTTTACATGGTAAATTTCATAATATAA MANGTTTAATGTCTGGVAATGGTGTAATTTACAAAAMAAGTCCACGTAGGCCAAAGATGG CTAANNCTGCATATAAGGVACGTGAATSCCAGTGGAAAGGTGTCTGAGGAGGGGCAGGGC CACAGGTGTCCTGACAGGGAACATCTTTGAAGGATCTGGMACAAACAAGGGCCCAGTTCA CAAACCACAGGTACACTCATTTTAGATAGGMCAGCAGAATAGGTGATGAAATTATACAGT TTYCACTTGTTGCCTACTTACTGAAGCAA

Entry Created: Aug 28 1995 Last Updated: Aug 28 1995

LIBRARY

Lib Name: Clontech human fetal brain polyA+ mRNA (#6535)

Organism: Homo sapiens

SUBMITTER

Name: Tsutomu Fujiwara

Lab: Otsuka GEN Research Institute Institution: Otsuka Pharmaceutical Co., Ltd

Address: 463-10 kagasuno Kawauchi-cho, Tokushima, Tokushima, 771-01

Japan

Tel: 0886-65-2888 Fax: 0886-37-1035

CITATIONS

Title: Fujiwara et al. (1995)

Authors: Fujiwara, T., Hirano, H., Katagiri, T., Kawai, A., Kuga, Y.,

Nagata, M., Okuno, S., Ozaki, K., Shimizu, F., Shimada, Y.,

Shinomiya, H., Takaichi, A., Takeda, S., Watanabe, T., Takahashi

,E., Hirai,Y., Maekawa,H., Shin,S., Nakamura,Y.

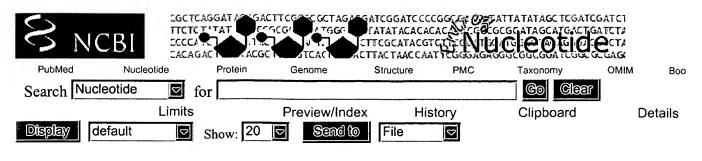
Year:

Status:

Unpublished

MAP DATA

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☐ 1: N45300. yz16g09.s1 Soares...[gi:1186466]

Links

### **IDENTIFIERS**

 dbEST Id:
 450829

 EST name:
 yz16g09.s1

 GenBank Acc:
 N45300

 GenBank gi:
 1186466

CLONE INFO

GDB Id:

Clone Id: IMAGE: 283264 (3')

Source: IMAGE Consortium, LLNL

3901792

DNA type: cDNA

PRIMERS

Sequencing: m13 -40 forward

PolyA Tail: Unknown

SEQUENCE

Quality:

High quality sequence stops at base: 386

Entry Created: Last Updated: Feb 13 1996 Feb 13 1996

COMMENTS

This clone is available royalty-free through LLNL ; contact the IMAGE Consortium ( $\underline{info@image.llnl.gov}$ ) for further

information.

LIBRARY

Lib Name: Soares multiple sclerosis 2NbHMSP

Organism: Homo sapiens

Sex: male

Tissue type: multiple sclerosis lesions

Develop. stage: Age 46

Lab host: DH10B (ampicillin resistant)

Vector: pT7T3D (Pharmacia) with a modified polylinker V TYPE:

phagemid

R. Site 1: Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer double-stranded cDNA was size selected, ligated to Eco RI adapters (Pharmacia), digested with Not I and cloned into the Not I and Eco RI sites of a modified pT7T3 vector

(Pharmacia). Library went through one round of normalization

to a Cot = 5. Library constructed by Bento Soares and

M.Fatima Bonaldo. RNA from 4 multiple sclerosis lesions from

one patient was kindly provided by Dr. Kevin G. Becker

(NINDS/NIH).

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax:

314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

The WashU-Merck EST Project

Authors:

Hillier, L., Clark, N., Dubuque, T., Elliston, K., Hawkins, M., Holman, M., Hultman, M., Kucaba, T., Le, M., Lennon, G., Marra, M. , Parsons, J., Rifkin, L., Rohlfing, T., Soares, M., Tan, F., Trevaskis, E., Waterston, R., Williamson, A., Wohldmann, P.,

Wilson, R.

Year:

1995

Status:

Unpublished

MAP DATA

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Genome

History

☐ 1: D80055. HUM015G10A Human ...[gi:1177932]

Limits

Links

### IDENTIFIERS

dbEST Id:

448123

EST name:

HUM015G10A

GenBank Acc: GenBank gi:

D80055

1177932

CLONE INFO

Clone Id:

GEN-015G10 (3')

DNA type:

**cDNA** 

PRIMERS

PolyA Tail:

Unknown

SEQUENCE

CCAAATTTCAAAAAGTTTTATTTTGAAAGAATGAGAGAAATAAAACAGAGAGGTATCAAT TACCAAGAACAATTACACTGAAGAAAACACAATAATAAGTACTCCCCACACAACCCCCC CCATTTCCCCATCCCTGGCACAATAATATTAAAACCACCAAAGCACCTAACAAGGAAA AACAACAGTACGTAATGAAAAAAGCAAATGTCCATACTGCTCAGTCCAACTAACCCTTAT GAAATGTCCTTCCCCCAGCTAAACCCTACCCACTGGAATGATAAAGAWATGTAGAGACAA CCCTAGGGGAGACTTGGAACTCTGCTTATACTAGCAAAGCTCAGTGAAGAATCAGTAAGA GTAGTGAATCTGTTTGGCAGTGAACACTGGATATAGCTTCTTTTTCAAATTTTGGATGAT TGCAGAGAACAGGTAGAGTTTGAGGCTCACAGACTTCTAACAGGACTGATCCCTGTTCCC

Entry Created: Last Updated:

Feb 9 1996 Feb 9 1996

LIBRARY

Lib Name:

Human fetal brain (TFujiwara)

TCAACCGTAACAGTGGGGBAGCTGCCAAATCCTGGGT

Organism:

Homo sapiens

Organ:

brain Develop. stage: fetal

SUBMITTER

Name:

Tsutomu Fujiwara

Lab: Institution: Otsuka GEN Research Institute Otsuka Pharmaceutical Co., Ltd

Address:

463-10 kagasuno Kawauchi-cho, Tokushima, Tokushima, 771-01

Japan

Tel: Fax: 0886-65-2888 0886-37-1035

CITATIONS

Title:

Fujiwara et al. (1995)

Authors:

Fujiwara, T., Hirano, H., Katagiri, T., Kawai, A., Kuga, Y., Nagata, M., Okuno, S., Ozaki, K., Shimizu, F., Shimada, Y.,

Shinomiya, H., Takaichi, A., Takeda, S., Watanabe, T., Takahashi

,E., Hirai,Y., Maekawa,H., Shin,S., Nakamura,Y.

Year:

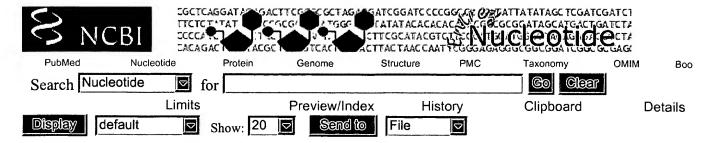
1995

Status:

Unpublished

MAP DATA

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□ 1: AA227221. zr20f08.s1 Strata...[gi:1848766]

Links

### **IDENTIFIERS**

dbEST Id:

866536

EST name:

zr20f08.s1

GenBank Acc:

AA227221

GenBank gi:

1848766

GDB Id:

5425825

# CLONE INFO

Clone Id:

IMAGE:663975 (3')

Source:

IMAGE Consortium, LLNL

Insert length:

1909

DNA type:

CDNA

#### PRIMERS

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

## SEQUENCE

CTGAATTAAAGAGATGAATTTATTGGTAAATAGATAAAATTAACACCTATTTTAATATAT CCAAACCCCTTCCTTATATATATTAGGTAAATTAAAAGAAAAAATTATCAAAGCAATAC ATATCAGGTGCTTCAGAGTTTTTCTCAAACAGTTACAAAAGAGATTAGGTTCCCAATCAG TTCACAAAAGCTAATTTAGAGAATGTAGCTTAACTACAGTACTGAGGTTGTCACACACTT AACTTTCGGTCTCTTGCTTATTTATTCATATCTGAGGTTCACTGTTTCTACTAGGATACA TTCCGCCCACACCCACACCTC

Entry Created:

Feb 19 1997

Last Updated:

Mar 11 1998

# COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

# LIBRARY

Lib Name:

Stratagene NT2 neuronal precursor 937230

Organism:

Homo sapiens

Organ:

brain

Tissue type:

neuroepithelial cells

Develop. stage: Ntera-2 neuroepithelial cells

Lab host:

SOLR (kanamycin resistant)

Vector:

pBluescript SK-

R. Site 1:

EcoRI

R. Site 2:

XhoI

Description:

Cloned unidirectionally. Primer: Oligo dT. Uninduced,

exponentially growing neuroepithelial cells (Ntera-2/c1.D1). Average insert size: 1.0 kb; Uni-ZAP XR Vector; ~5' adaptor

sequence: 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5'

CTCGAGTTTTTTTTTTTTTTT 3'

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax:

314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

WashU-NCI human EST Project

Authors:

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost ,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra ,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan ,F., Theising,B., White,Y., Wylie,T., Waterston,R., Wilson

,R.

Year:

1997

Status:

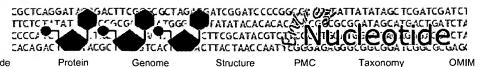
Unpublished

## MAP DATA

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Taxonomy





Structure

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Display Show: 20 default  $\overline{\mathbf{v}}$ <u>ග් bm</u>ෙ File  $\Diamond$ 

☐ 1: N64489. yz90e07.s1 Soares...[gi:1212318]

Links

Details

Boo

### **IDENTIFIERS**

dbEST Id:

470240

EST name:

yz90e07.s1

GenBank Acc: GenBank gi:

N64489 1212318

GDB Id:

3906180

CLONE INFO

Clone Id:

IMAGE:290340 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

**CDNA** 

PRIMERS

Sequencing:

m13 -40 forward

PolyA Tail:

Unknown

SEQUENCE

GATTAACTATGTGACTAAATTATATTCAAATTTTATGAACAGAAAATGATAAAATGTTA TCAGCTAATAAAGAGATTATCAAAGAGTAAGCAACCAAAACAAGTAGGCAAAAAGCATCA GAGAGTAATTAATACAAAGATGATGTTGTTTTTCTGGATTTCATAATGTTTATCATAGTT GTCAACTTTTCTCATTCAAAAAAACCCTTATTTTTATACCTAATTTTAATTAAAAATTTT TCAGTTTGTATTAAAGAGGACTCCCCAAATTATATGAGTTTCCAACTTCATAAAACCTAA ATCTGTCTTTGTTCATATCAGATAAAAATAGGCCACACAGACTGCCAAGTAGGTACAGTC TTGGAACTGTCTGTGGTGCTGGACCCAAGGTTCACTTGGGCTCTCTCCATGGGTACTTAC TGGCCCAAGCCAAAGCTG

Quality:

High quality sequence stops at base: 416

Entry Created: Last Updated:

Mar 1 1996 Mar 1 1996

COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

LIBRARY

Lib Name:

Soares multiple sclerosis 2NbHMSP

Organism:

Homo sapiens

Sex:

male

Tissue type:

multiple sclerosis lesions

Develop. stage: Age 46

Lab host:

DH10B (ampicillin resistant)

Vector:

pT7T3D (Pharmacia) with a modified polylinker V\_TYPE:

phagemid

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

1st strand cDNA was primed with a Not I - oligo(dT) primer

## SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel:

314 286 1800 314 286 1810

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Parsons,J., Rifkin,L., Rohlfing,T., Soares,M., Tan,F.,
Trevaskis,E., Waterston,R., Williamson,A., Wohldmann,P.,

Wilson, R.

Year:

1995

Status:

Unpublished

## MAP DATA

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GATCGGATCCCCGGCER (2) TATTATATAGCTCGATCGATCT FATATACACACACA (2) CGCGCGGTAGCATGACTGATCTA CGCTCAGGATA CGCTAGA A TIGGE CTTCGCATACGTCT CACAGACT ACTTACTAACCAATTEGGG.

Structure

**PMC** 

Taxonomy

Genome Search Nucleotide Clear  $\triangle$ **(490** for Clipboard

Limits History Preview/Index Show: 20 Display default  $\nabla$ od bnes File  $\triangle$ 

Protein

☐ 1: AA478962. zv18d05.s1 Soares...[gi:2207596]

Nucleotide

Links

Details

Boo

### IDENTIFIERS

PubMed

dbEST Id:

1124075

EST name:

zv18d05.s1 AA478962

GenBank Acc:

2207596

GenBank gi: GDB Id:

5976866

# CLONE INFO

Clone Id:

IMAGE: 753993 (3')

Source:

IMAGE Consortium, LLNL

DNA type:

**CDNA** 

## PRIMERS

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

## SEQUENCE

TTTTTTTTTTTTTTTGGTCATACTTCACTTTATTATTATTATTATCATA CATGGTTACTATTCCAATCTTTCATGCAGACAAAATAAACAATATAAAATACATAATGC ACTTTGATAATTTTAACCATACATAAAATATGGAGTAATGGAAGCTATGTTACATGGATA TTTTACAAAGGAAAAAAGATGACTTTTATAATAACACATCCAGATGAAATTTATCATTA AATTTTGGATTTCATATGATGTTAAGTATGGATATATTCAAAACAATTACTATTTATAGA ACCAATTTGATATTTTGTCATTTAAAATAATGAATACTATGTAAATGAGTACTTATAAAA

ATATTTTTAGGCAAAAAG

Quality:

High quality sequence stops at base: 249

Entry Created:

Jun 19 1997

Last Updated:

Aug 8 1997

## COMMENTS

This clone is available royalty-free through LLNL; contact the IMAGE Consortium (info@image.llnl.gov) for further

information.

# LIBRARY

Lib Name:

Soares\_NhHMPu\_S1 Homo sapiens

Organism:

mixed (see below)

Organ: Tissue type:

Pooled human melanocyte, fetal heart, and pregnant uterus

Lab host:

Vector:

pT7T3D-Pac (Pharmacia) with a modified polylinker

R. Site 1:

Not I

R. Site 2:

Eco RI

Description:

Equal amounts of plasmid DNA from three normalized libraries (melanocyte 2NbHM, pregnant uterus NbHPU, and fetal heart NbHH19W) were mixed, and ss circles were made in vitro. Following HAP purification, this DNA was used as tracer in a

subtractive hybridization reaction. The driver was

PCR-amplified cDNAs from pools of 5,000 clones made from the same 3 libraries. The pools consisted of I.M.A.G.E. clones

260232-265223, 340488-345479, and 484488-489479.

SUBMITTER

Name:

Wilson RK

Institution:

Washington University School of Medicine

Address:

4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: Fax: 314 286 1800 314 286 1810

E-mail:

est@watson.wustl.edu

CITATIONS

Title:

WashU-Merck EST Project 1997

Authors:

Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost, S., Kucaba, T., Lacy, M., Le, N., Lennon, G., Marra, M., Martin, J., Moore, B., Schellenberg, K., Steptoe, M., Tan, F., Theising, B., White, Y., Wylie, T., Waterston, R., Wilson, R.

Year:

1997

Status:

Unpublished

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☐ 1: AA485973. ab11e10.s1 Strata...[gi:2215124]

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### IDENTIFIERS

dbEST Id:

1130535

EST name: GenBank Acc: ab11e10.s1 AA485973

GenBank gi:

2215124

# CLONE INFO

Clone Id:

IMAGE:840522 (3')

Source:

IMAGE Consortium, LLNL

Insert length:

967 **CDNA** 

DNA type:

## **PRIMERS**

Sequencing:

-41m13 fwd. ET from Amersham

PolyA Tail:

Unknown

## SEQUENCE

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TTTGAAATAATATTCTCCCCA

Quality:

High quality sequence stops at base: 262

Entry Created:

May 29 1997

Last Updated:

Mar 5 1998

## COMMENTS

This clone is available royalty-free through LLNL; contact

the IMAGE Consortium (info@image.llnl.gov) for further

information.

PUTATIVE ID

Assigned by submitter

contains MER30.t3 MER30 repetitive element;

# LIBRARY

Lib Name:

Stratagene lung (#937210)

Organism:

Homo sapiens

Sex: Organ: male

Develop. stage: 72 years

lung

Lab host:

Vector:

SOLR cells (kanamycin resistant)

R. Site 1:

pBluescript SK-EcoRI

R. Site 2:

XhoI

Description: Cloned unidirectionally. Primer: Oligo dT. normal lung.

Average insert size: 1.0 kb; Uni-ZAP XR Vector; ~5' adaptor

sequence: 5' GAATTCGGCACGAG 3' ~3' adaptor sequence: 5'

CTCGAGTTTTTTTTTTTTTTTTT 3'

SUBMITTER

Name: Wilson RK

Institution: Washington University School of Medicine

Address: 4444 Forest Park Parkway, Box 8501, St. Louis, MO 63108

Tel: 314 286 1800 Fax: 314 286 1810

E-mail: est@watson.wustl.edu

CITATIONS

Title: WashU-NCI human EST Project

Authors: Hillier, L., Allen, M., Bowles, L., Dubuque, T., Geisel, G., Jost

,S., Krizman,D., Kucaba,T., Lacy,M., Le,N., Lennon,G., Marra,M., Martin,J., Moore,B., Schellenberg,K., Steptoe,M., Tan,F., Theising,B., White,Y., Wylie,T., Waterston,R., Wilson

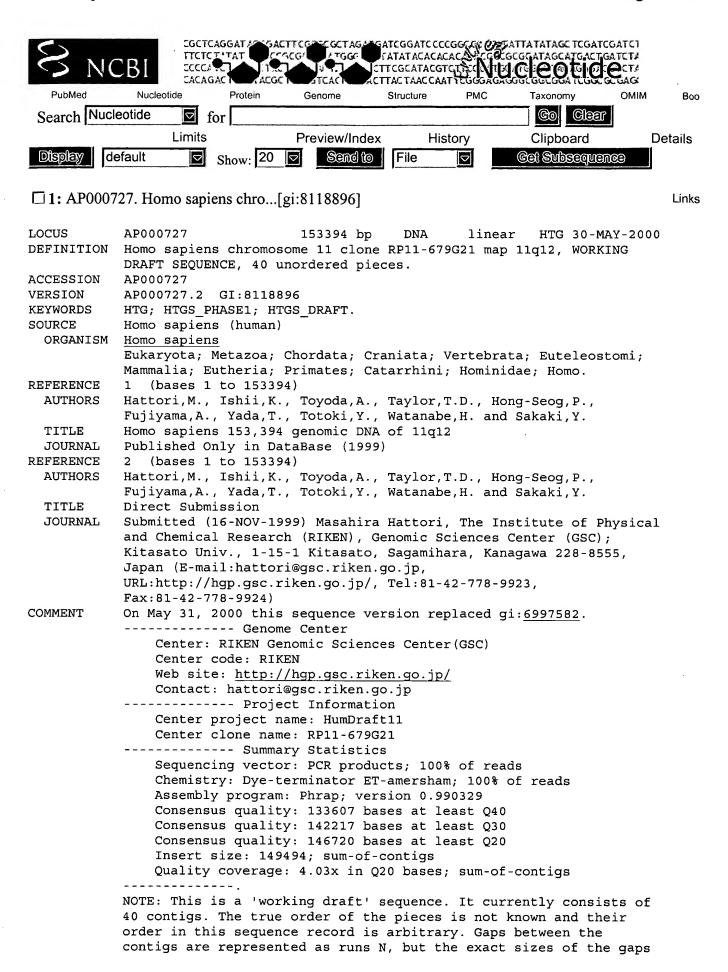
,R.

Year: 1997

Status: Unpublished

MAP DATA

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are unknown. This record will be updated with the finished sequence as soon as it is available and the accession number will be preserved

1 11871 contig of 11871 bp in length
11972 22347 contig of 10376 bp in length
22448 33540 contig of 11093 bp in length
33641 41577 contig of 7937 bp in length
41678 45975 contig of 4298 bp in length
46076 53127 contig of 7052 bp in length

46076 53127 contig of 7052 bp in length 53228 60314 contig of 7087 bp in length 60415 65544 contig of 5130 bp in length 71027 contig of 65645 5383 bp in length 75422 contig of 4295 bp in length 71128 75523 3752 bp in length 79274 contig of 79375 84398 contig of 5024 bp in length 84499 88990 contig of 4492 bp in length 89091 93436 contig of 4346 bp in length 96655 contig of 3119 bp in length 93537 96756 99175 contig of 2420 bp in length 99276 102659 contig of 3384 bp in length 102760 105824 contig of 3065 bp in length 105925 109361 contig of 3437 bp in length 109462 112303 contig of 2842 bp in length 115484 contig of 3081 bp in length 112404 115585 118972 contig of 3388 bp in length 119073 121620 contig of 2548 bp in length 121721 124322 contig of 2602 bp in length 124423 127088 contig of 2666 bp in length 127189 130306 contig of 3118 bp in length 130407 131809 contig of 131910 133740 contig of 133841 136109 contig of 136210 137539 contig of 1403 bp in length 1831 bp in length 2269 bp in length 1330 bp in length 137640 139390 contig of 1751 bp in length 139491 140939 contig of 1449 bp in length 141040 142493 contig of 1454 bp in length 142594 144234 contig of 1641 bp in length 144335 145850 contig of 1516 bp in length 145951 147405 contig of 1455 bp in length 147506 149002 contig of 1497 bp in length 149103 150523 contig of 1421 bp in length 150624 152235 contig of 1612 bp in length 153394 contig of 1059 bp in length 152336

Sequence updated (26-May-2000).

\* NOTE: This is a 'working draft' sequence. It currently

\* consists of 40 contigs. The true order of the pieces

\* is not known and their order in this sequence record is

\* arbitrary. Gaps between the contigs are represented as

\* runs of N, but the exact sizes of the gaps are unknown.

\* This record will be updated with the finished sequence

\* as soon as it is available and the accession number will

\* be preserved.

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\* 33641 41577: contig of 7937 bp in length

\* 41578 41677: gap of 100 bp

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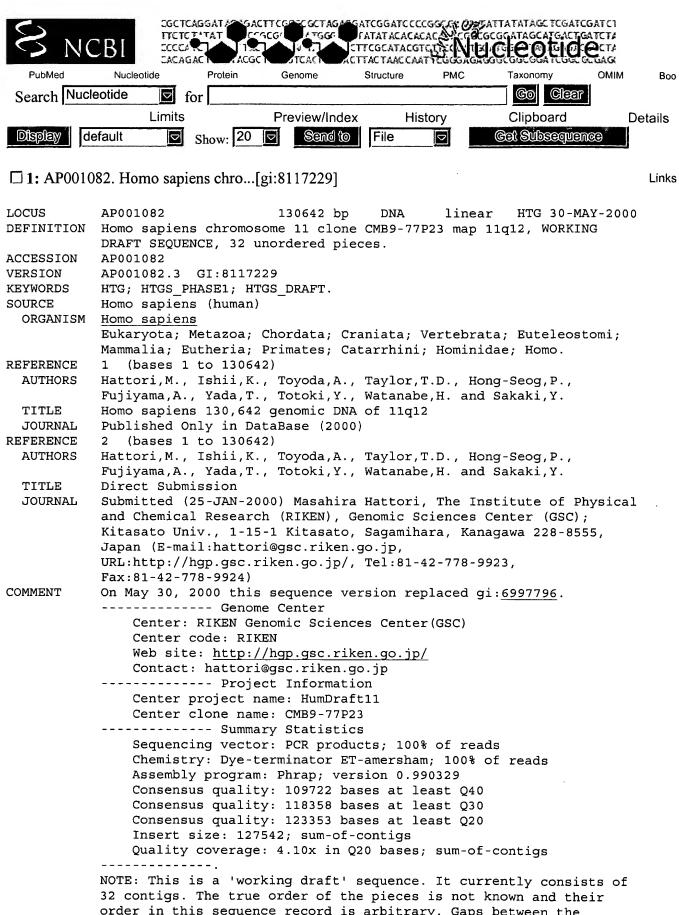
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Jul 8 2003 12:22:35



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